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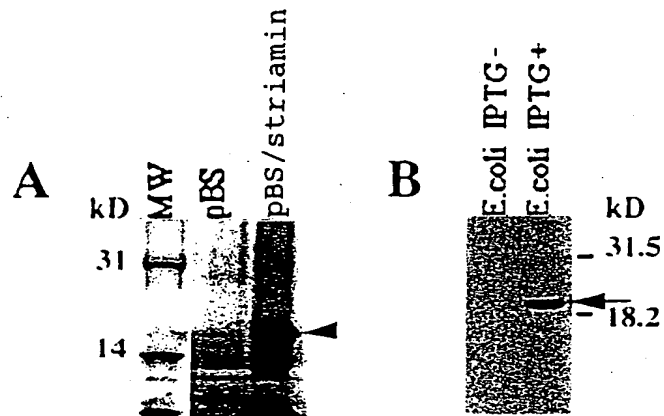
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FIG. 1

5' -GCAGGTCTGAGTTCAAGGACAGCCTGGTCTACGCATTGAGTTCTAGAACAGCCAAGGCTACACAAAGA																										68	
ATCCCTGTCTTAAAAAACAAAACAAATAAAAAACAAACAACAAAAACAAAACAAAACAAAACAAAGACCA																										147	
ATGGGGGAAAAAAGAAAGAAAAACAAGAAAAGAAAAAGAATAGCTTCCTCGTTCTCTGAGGGTAGTTTTAGTAATG																										226	
AATGCTCAAAGCTCCACAGTCTATGGCACCCAAGTGGTATTCTTATATTGTTTGCAACTAACTATCC ATG AAA GGC																										302	
																										M K G	3
<u>CTG GCT GGC GAG TGG CAT CAG GAC TCT GGC CTA GAC ATC AGG GAG AAG GCA GAA GAC TTC</u>																										362	
<u>L A G E W H Q D S G L D I R E K A E D F</u>																										23	
<u>TCC CTG CCC TGG CTG CTG CCT AGA TTG ATG GCC TTA GTC ATG CAG GAA GAA GGA AGG TTC</u>																										422	
<u>S L P W L L P R L M A L V M Q E E G R F</u>																										43	
<u>AGA AGT GAC AGG AAT CAT GGG TAT TTA AGG GAA TGG CTT AGG ATT CAG GCA CTG ACA GCT</u>																										482	
<u>R S D R N H G Y L R E W L R I Q A L T A</u>																										163	
<u>TGT CTG CCT TCC CCT CTG GGG AGG GTC CAC TAT GCC CAG TGT TCA CCG AAA CAA AAA GGA</u>																										542	
<u>C L P S P L G R V H Y A Q C S P K Q K G</u>																										83	
<u>AGG CTG CCA AGA GGC TGG GCT TCT CTG CCA TCC CTA AGT GTG CTT GTC AGG GCT CTG AGA</u>																										602	
<u>R L P R G W A S L P S L S V L V R A L R</u>																										103	
<u>GCG TCT AAC GCT TTC TCA CTC GGG AAC TAC TAC TGC TGT CCG TGG AGG GGG ACA AGA TGG</u>																										662	
<u>A S N A F S L G N Y Y C C P W R G T R W</u>																										123	
<u>GCC AAA GGG CAG CCG GGA GAG TGG GCA AGG CCA AGG GCG AGC TCA CCT TCT GCG CAC CGA</u>																										722	
<u>A K G Q P G E W A R P R A S S P S A H R</u>																										143	
<u>GAA GGC GTG GCA GTG ACA TGA AAC ATT AGG TCA CAT GGC CTT CCC CTC CGG CCT TAG CGT</u>																										782	
<u>E G V A V T *</u>																										149	
GCCTACACATCTGCACAGAGAAGGAGAAGAGGTTGAGGAGAGAATAGAGCAGTACAGAGCCAACTCAAACAGCAGCAGC																										861	
AGACAGAGAGCAGCAGACAGAAAGGTGAGGGCGCAGGAGGAGAGGAGCGGGCCAGAAGGGACAGTGAGCAGACAGGGAAA																										940	
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T-3'																										2679	

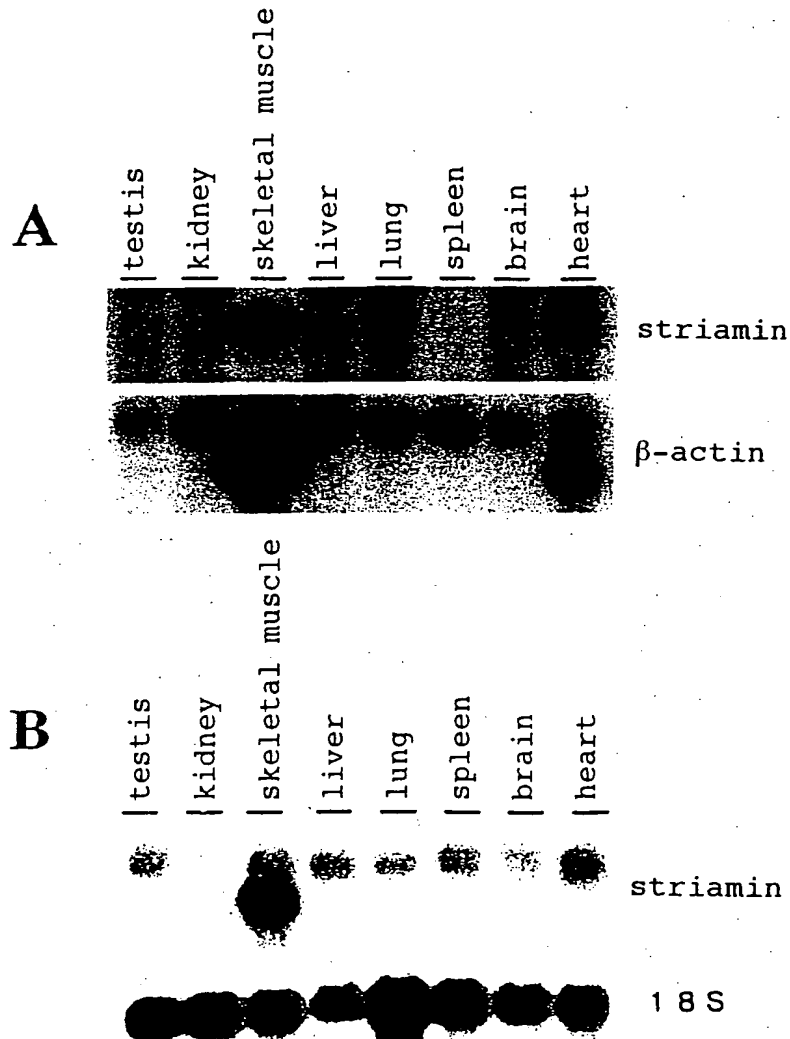
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Figure 2



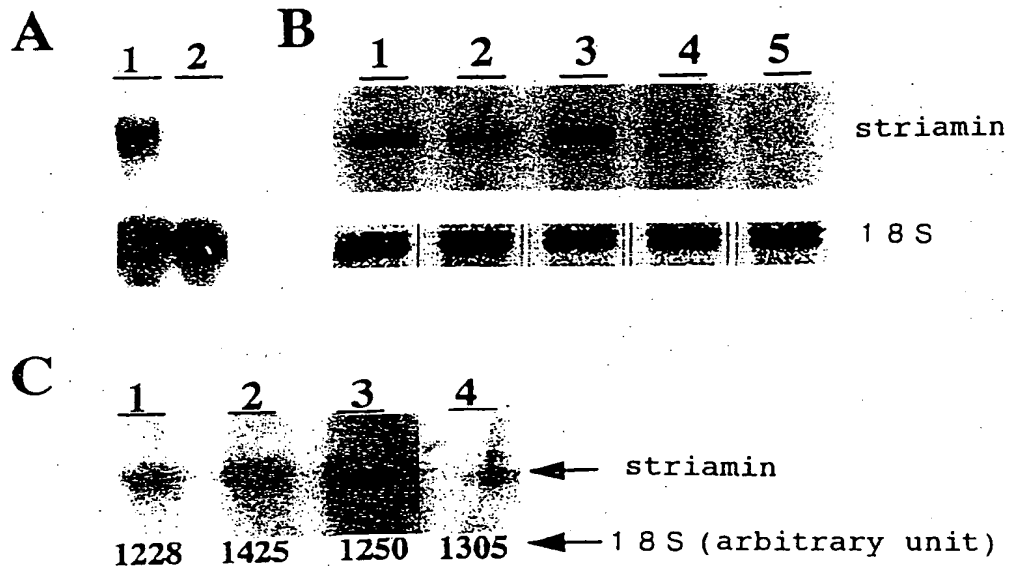
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Figure 3



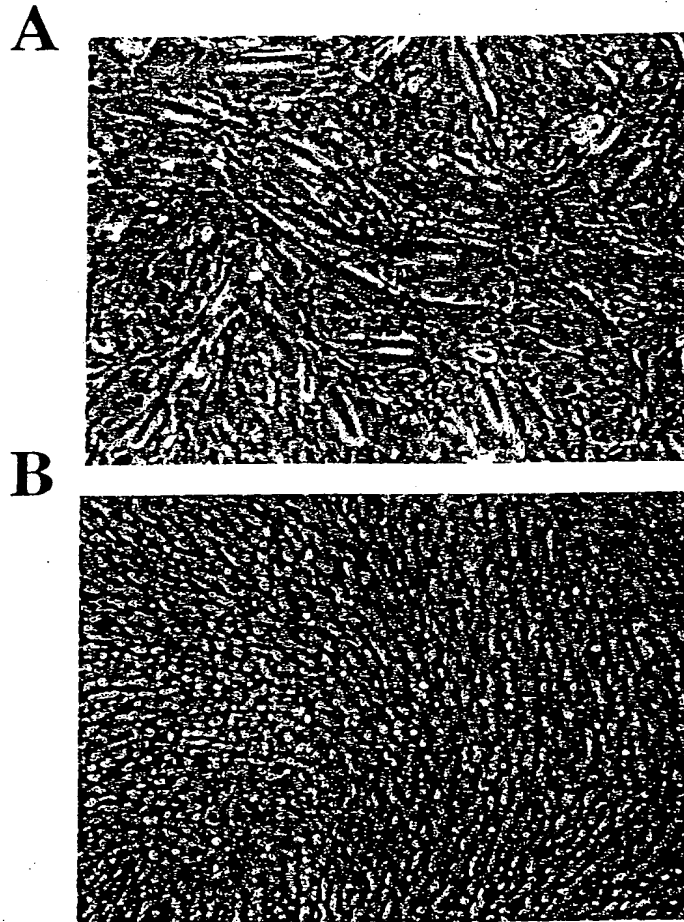
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Figure 4



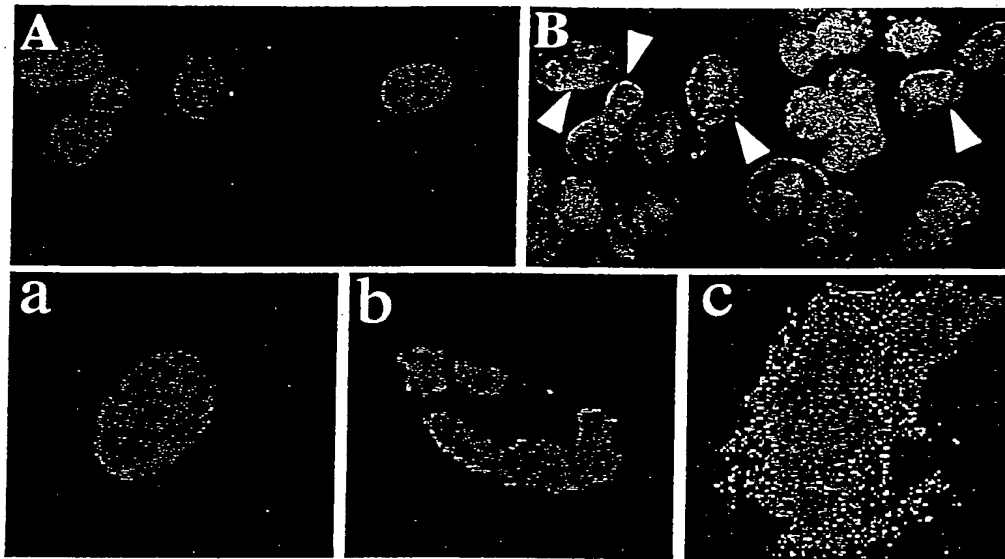
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Figure 5



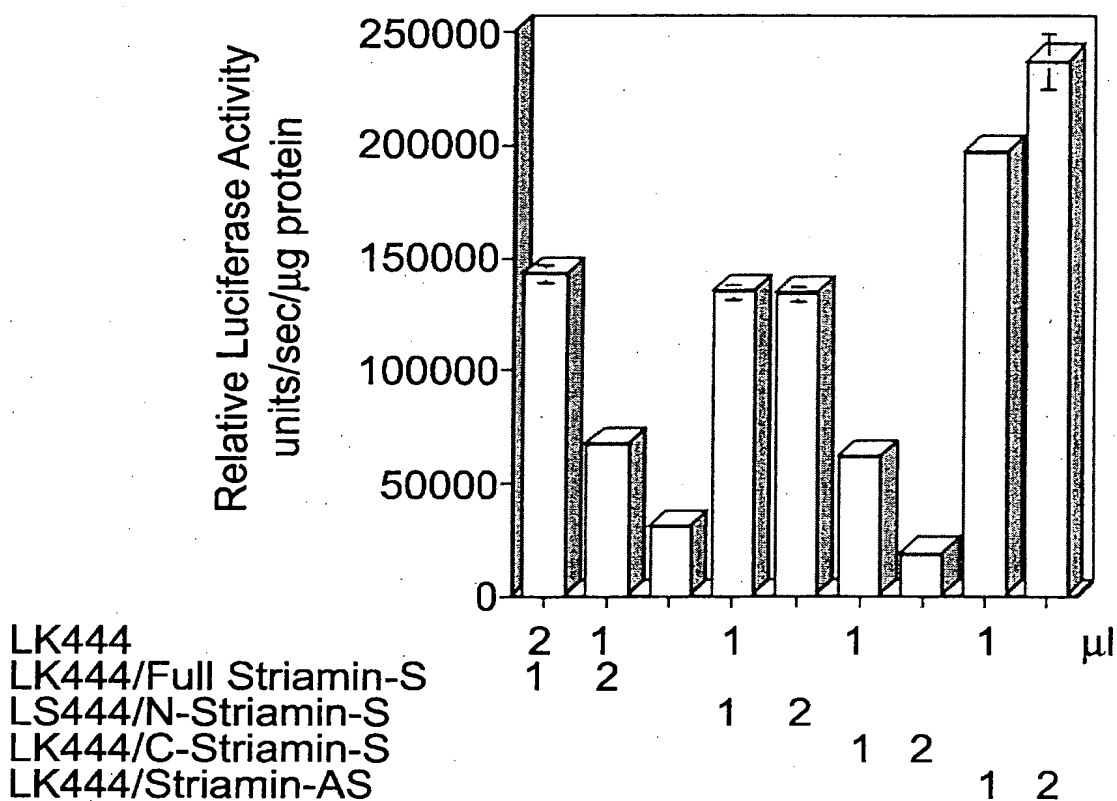
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Figure 6



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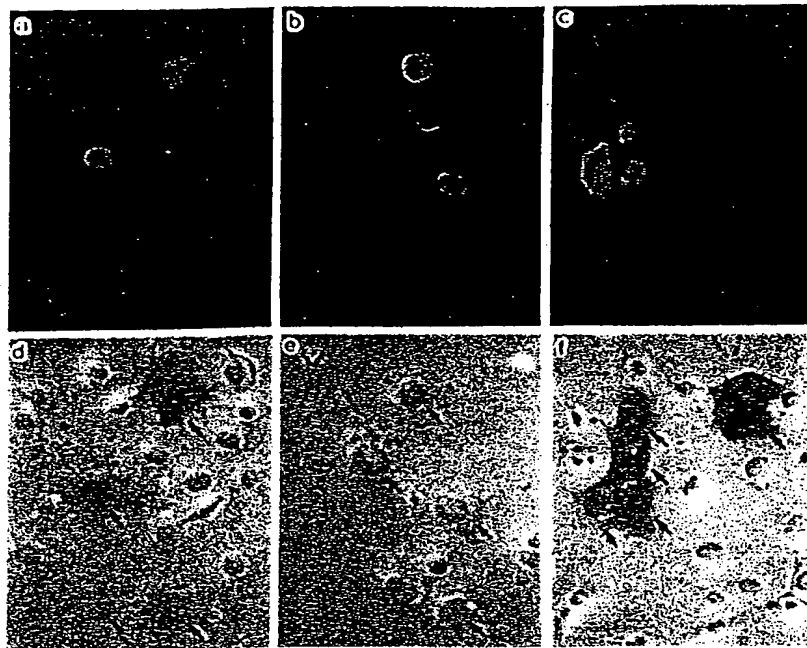
FIG. 7





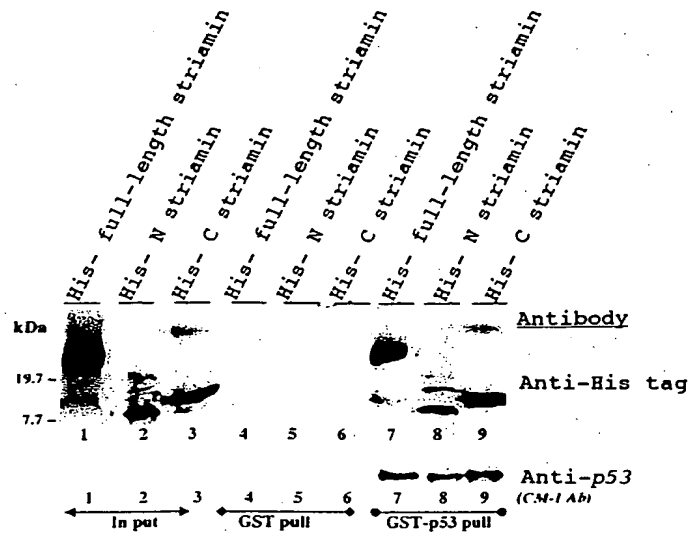
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Figure 8



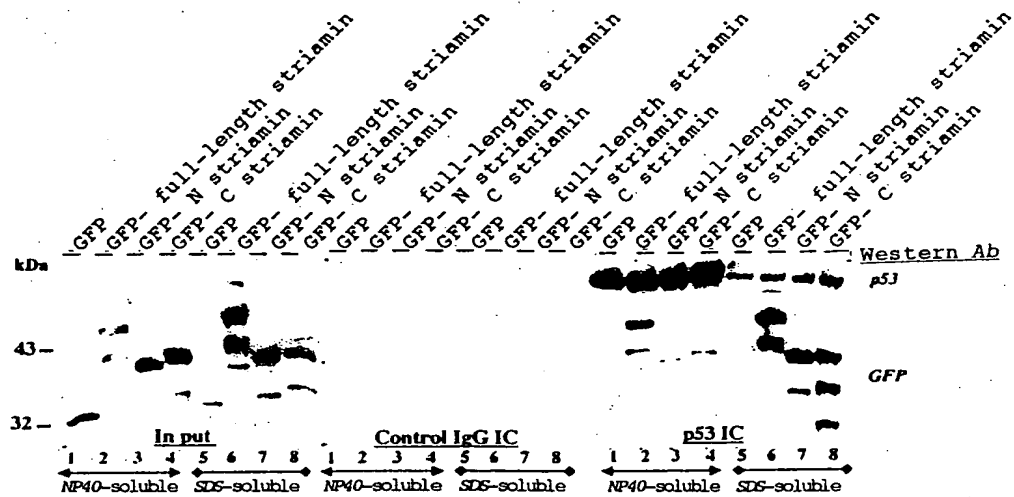
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Figure 9



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Figure 10



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FIG. 11

